

FIGURE 1

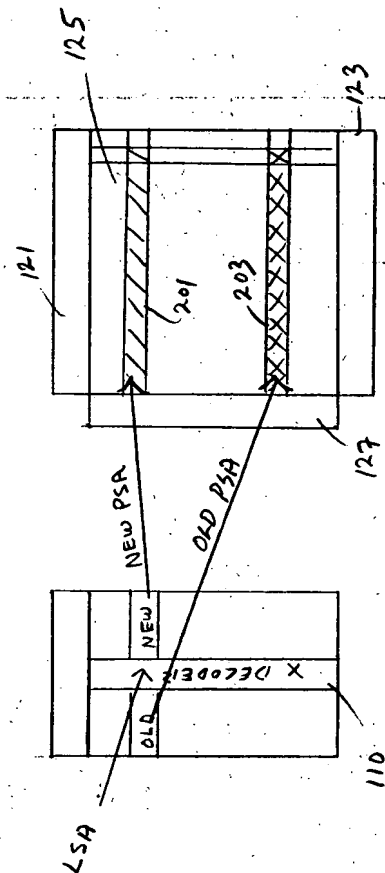


FIGURE 2

LSA 110

1	
2	
3	
4	
5	
6	
7	
X	

301 303 305

PSA 125 321 323

1	
2	
3	
4	
5	
6	
7	
8	

127

FIGURE 3a

110 125 321 323

1	A	A	A	A	A	A	A	A
2	B	B	B	B	B	B	B	B
3	X	X	X	X	X	X	X	X
4								
5	C	C	C	C	C	C	C	C
6	X	X	X	X	X	X	X	X
7	E	E	E	E	E	E	E	E
8	X	X	X	X	X	X	X	X

127

FIGURE 3d

LSA 110

1	
2	
3	5
4	
5	
6	
7	
X	

301 303 305

PSA 125 321 323

1	A	A	A	A	A	A	A	A
2	B	B	B	B	B	B	B	B
3	X	X	X	X	X	X	X	X
4								
5	C	C	C	C	C	C	C	C
6								
7								
8								

127

FIGURE 3b

110 125 321 323

1	A	A	A	A	A	A	A	A
2	B	B	B	B	B	B	B	B
3								
4								
5	C	C	C	C	C	C	C	C
6								
7	E	E	E	E	E	E	E	E
8								

127

FIGURE 3e

110

1	
2	
3	5
4	
5	
6	
7	
X	

301 303 305

125 321 323

1	A	A	A	A	A	A	A	A
2	B	B	B	B	B	B	B	B
3	X	X	X	X	X	X	X	X
4								
5	C	C	C	C	C	C	C	C
6	D	D	D	D	D	D	D	D
7	E	E	E	E	E	E	E	E
8	F	F	F	F	F	F	F	F

127

FIGURE 3c

110 125 321 323

1	A	A	A	A	A	A	A	A
2	B	B	B	B	B	B	B	B
3	G	G	G	G	G	G	G	G
4								
5	C	C	C	C	C	C	C	C
6								
7	E	E	E	E	E	E	E	E
8								

127

FIGURE 3f

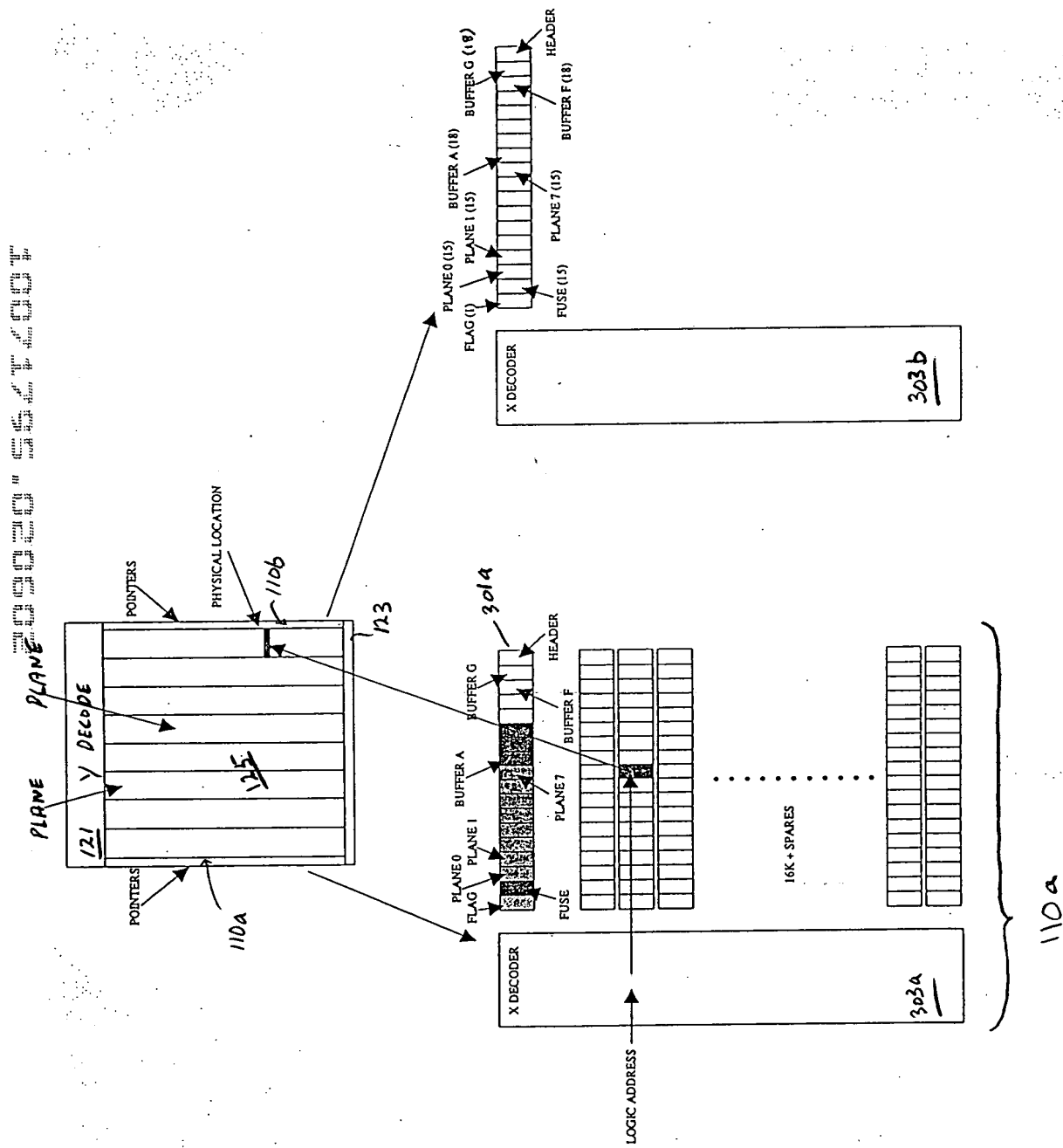


FIGURE 4